

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
3 November 2005 (03.11.2005)

PCT

(10) International Publication Number
WO 2005/103239 A3

(51) International Patent Classification:
C12N 9/04 (2006.01) **C12N 15/31** (2006.01)

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(21) International Application Number:
PCT/EP2005/002557

(22) International Filing Date: 10 March 2005 (10.03.2005)

(25) Filing Language: English

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(26) Publication Language: English

(81) Designated States (*unless otherwise indicated, for every kind of national protection available*): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BW, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NA, NI, NO, NZ, OM, PG, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, SM, SY, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(30) Priority Data:
10 2004 014 274.2 22 March 2004 (22.03.2004) DE

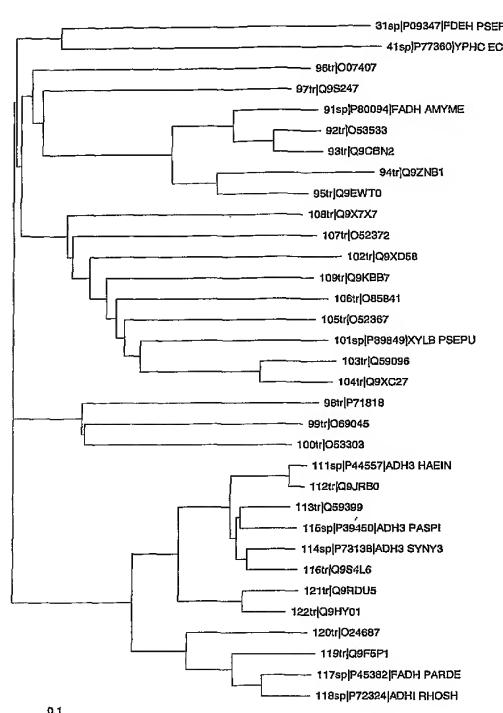
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[Continued on next page]

(54) Title: NOVEL ALCOHOL DEHYDROGENASES



Overview of cluster 2 (= primer group 2), based on 33 sequences

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(84) **Designated States** (*unless otherwise indicated, for every kind of regional protection available*): European (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IS, IT, LT, LU, MC, NL, PL, PT, RO, SE, SI, SK, TR).

— *with amended claims*

(88) **Date of publication of the international search report:**
9 March 2006

Declaration under Rule 4.17:
— *of inventorship (Rule 4.17(iv))*

Date of publication of the amended claims: 27 April 2006

Published:
— *with international search report*

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

AMENDED CLAIMS

[received by the International Bureau on 27 January 2006 (27.01.06);
original claim 1 replaced by new claims 1; remaining claims unchanged (3 pages)]

1. A polypeptide which has the biological activity of
an NAD- or NADP-dependent alcohol dehydrogenase
5 and which comprises or has one of the following
sequences: the sequence of SEQ ID NO.: 1, the
sequence of SEQ ID NO.: 2, the sequence of SEQ ID
NO.: 3 or a sequence which is at least 90%
identical to the sequence of SEQ ID NO.: 3, the
10 sequence of SEQ ID NO.: 4, the sequence of SEQ ID
NO.: 5 or a sequence which is at least 90%
identical to the sequence of SEQ ID NO.: 5, the
sequence of SEQ ID NO.: 6 or a sequence which is
at least 90% identical to the sequence of SEQ ID
15 NO.: 6, the sequence of SEQ ID NO.: 7 or a
sequence which is at least 70% identical to the
sequence of SEQ ID NO.: 7, the sequence of SEQ ID
NO.: 8 or a sequence which is at least 70%
identical to the sequence of SEQ ID NO.: 8, the
20 sequence of SEQ ID NO.: 9 or a sequence which is
at least 70% identical to the sequence of SEQ ID
NO.: 9, the sequence of SEQ ID NO.: 10 or a
sequence which is at least 70% identical to the
sequence of SEQ ID NO.: 10, the sequence of SEQ ID
25 NO.: 11 or a sequence which is at least 70%
identical to the sequence of SEQ ID NO.: 11, the
sequence of SEQ ID NO.: 12 or a sequence which is
at least 60% identical to the sequence of SEQ ID
NO.: 12, the sequence of SEQ ID NO.: 13 or a
30 sequence which is at least 60% identical to the
sequence of SEQ ID NO.: 13, the sequence of SEQ ID
NO.: 14 or a sequence which is at least 75%
identical to the sequence of SEQ ID NO.: 14, the
sequence of SEQ ID NO.: 15 or a sequence which is
35 at least 95% identical to the sequence of SEQ ID
NO.: 15, the sequence of SEQ ID NO.: 16 or a
sequence which is at least 95% identical to the

sequence of SEQ ID NO.: 16, the sequence of SEQ ID NO.: 17 or a sequence which is at least 75% identical to the sequence of SEQ ID NO.: 17, the sequence of SEQ ID NO.: 18 or a sequence which is at least 70% identical to the sequence of SEQ ID NO.: 18, the sequence of SEQ ID NO.: 19 or a sequence which is at least 70% identical to the sequence of SEQ ID NO.: 19, the sequence of SEQ ID NO.: 20 or a sequence which is at least 60%
5 identical to the sequence of SEQ ID NO.: 20, the sequence of SEQ ID NO.: 21 or a sequence which is at least 90% identical to the sequence of SEQ ID NO.: 21, the sequence of SEQ ID NO.: 22 or a sequence which is at least 70% identical to the
10 sequence of SEQ ID NO.: 22, the sequence of SEQ ID NO.: 23 or a sequence which is at least 55% identical to the sequence of SEQ ID NO.: 23, the sequence of SEQ ID NO.: 24 or a sequence which is at least 65% identical to the sequence of SEQ ID
15 NO.: 24, the sequence of SEQ ID NO.: 25 or a sequence which is at least 55% identical to the sequence of SEQ ID NO.: 25, the sequence of SEQ ID NO.: 26 or a sequence which is at least 55% identical to the sequence of SEQ ID NO.: 26, the sequence of SEQ ID NO.: 27 or a sequence which is at least 55% identical to the sequence of SEQ ID NO.: 27, the sequence of SEQ ID NO.: 28 or a sequence which is at least 75% identical to the sequence of SEQ ID NO.: 28, the sequence of SEQ ID
20 NO.: 29 or a sequence which is at least 70% identical to the sequence of SEQ ID NO.: 29, the sequence of SEQ ID NO.: 30 or a sequence which is at least 60% identical to the sequence of SEQ ID NO.: 30, the sequence of SEQ ID NO.: 31 or a sequence which is at least 55% identical to the sequence of SEQ ID NO.: 31, the sequence of SEQ ID
25 NO.: 32 or a sequence which is at least 55%

identical to the sequence of SEQ ID NO.: 32, the sequence of SEQ ID NO.: 33 or the sequence of SEQ ID NO.: 34.

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